

Figure 12 is a chart that depicts results of microarray expression analysis of the ovarian carcinoma sequence designated 8e.

Figure 13 is a chart that depicts results of microarray expression analysis of the ovarian carcinoma sequence designated 12c.

Figure 14 is a chart that depicts results of microarray expression analysis of the ovarian carcinoma sequence designated 12h.

At page 48, immediately preceding the table on this page, please insert the following label:

TABLE 1 – Ovarian Carcinoma cDNAs Identified by Microarray Analysis.

*In the Claims:*

Please cancel claims ~~1, 2, 9, 12, 21, 23, 24, and 69.~~

Please add claims 73-81.

73. (New) An isolated polypeptide comprising SEQ ID NO:392.

74. (New) An isolated polypeptide comprising an amino acid sequence having at least 90% identity to SEQ ID NO:392, wherein said polypeptide is over expressed in ovarian cancer relative to normal ovarian tissue, and wherein said polypeptide is not SEQ ID NO:393.

75. (New) An isolated polypeptide comprising amino acid residues 1-27 of SEQ ID NO:392, wherein said polypeptide is over expressed in ovarian cancer relative to normal ovarian tissue.

76. (New) An isolated polypeptide comprising amino acid residues 1-27 of SEQ ID NO:392, or a sequence having at least 90% identity to said polypeptide, wherein said polypeptide is over expressed in ovarian cancer relative to normal ovarian tissue.

77. (New) An isolated polypeptide comprising an amino acid sequence having at least 90% identity to amino acid residues 1-27 of SEQ ID NO:392,